

ID AC005973 standard; DNA; HTG; 150587 BP.
 XX
 AC AC005973;
 XX
 SV AC005973.6
 XX
 DT 19-NOV-1998 (Rel. 57, Created)
 DT 11-AUG-2000 (Rel. 64, Last updated, Version 25)
 XX
 DE Homo sapiens chromosome 17 clone RP11-177H5 map 17, *** SEQUENCING IN
 DE PROGRESS ***, 2 ordered pieces.
 XX
 KW HTG; HTGS_PHASE2.
 XX
 OS Homo sapiens (human)
 OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia;
 OC Eutheria; Primates; Catarrhini; Hominidae; Homo.
 XX
 RN [1]
 RP 1-150587
 RA Birren B., Linton L., Nusbaum C., Lander E.;
 RT "Homo sapiens chromosome 17, clone RP11-177H5";
 RL Unpublished.
 XX
 RN [2]
 RP 1-150587
 RA Birren B., Linton L., Nusbaum C., Lander E., Allen N., Anderson M.,
 RA Baker J., Baldwin J., Barna N., Beckerly R., Benn J., Boutwell C.,
 RA Brown A., Castle A., Cerny J., Colangelo M., Collins S., Collymore A.,
 RA Cooke P., Corliss D., Depayre E., Devon K., Dewar K., Donelan L.,
 RA Ferreira P., FitzHugh W., Forrest C., Funke R., Gage D., Gardyna S.,
 RA Geraigery K., Grant G., Hagos B., Heaford A., Herena L., Horton L.,
 RA Howland J.C., Jacotot L., Jones C., Kann L., Karatas A., Lehoczký J.,
 RA Macdonald P., Marquis N., McEwan P., McGurk A., McKernan K., Meldrim J.,
 RA Molla M., Morris W., Morrow J., Mychaleckyj J., Nahf R., Naylor J.,
 RA Niloff M., O'Connor T., O'Donnell P., Pavlin B., Peterson K., Riley R.,
 RA Roberts D., Roy A., Severy P., Stange-Thomann N., Stilwell J.,
 RA Stojanovic N., Stone C., Subramanian A., Tesfaye S., Tichovolsky N.,
 RA Torruella-Miller I., Vassiliev H., Vo A., Wagner A., Wheeler J., Wu Y.,
 RA Wyman D., Ye W.J., Zhao J., Zody M.;
 RT ;
 RL Submitted (18-NOV-1998) to the EMBL/GenBank/DDBJ databases.
 RL Whitehead Institute/MIT Center for Genome Research, 320 Charles Street,
 RL Cambridge, MA 02141, USA
 XX
 CC On Aug 10, 2000 this sequence version replaced gi:8516060.
 CC All repeats were identified using RepeatMasker:
 CC Smit, A.F.A. & Green, P. (1996-1997)
 CC <http://ftp.genome.washington.edu/RM/RepeatMasker.html>
 CC ----- Genome Center
 CC Center: Whitehead Institute/ MIT Center for Genome Research
 CC Center code: WIBR
 CC Web site: <http://www-seq.wi.mit.edu>
 CC Contact: sequence_submissions@genome.wi.mit.edu
 CC ----- Project Information
 CC Center project name: L485
 CC Center clone name: 177_H_5
 CC -----
 CC * NOTE: This is a 'working draft' sequence. It currently
 CC * consists of 2 contigs. Gaps between the contigs
 CC * are represented as runs of N. The order of the pieces
 CC * is believed to be correct as given, however the sizes
 CC * of the gaps between them are based on estimates that have
 CC * provided by the submittor.

CC * This sequence will be replaced
 CC * by the finished sequence as soon as it is available and
 CC * the accession number will be preserved.
 CC * 1 76468: contig of 76468 bp in length
 CC * 76469 76568: gap of 100 bp
 CC * 76569 150587: contig of 74019 bp in length.

XX

FH Key Location/Qualifiers

FH

FT source 1..150587

FT /chromosome="17"

FT /db_xref="taxon:9606"

FT /organism="Homo sapiens"

FT /map="17"

FT /clone="RP11-177H5"

FT /clone_lib="RPCI-11 Human Male BAC"

XX

SQ Sequence 150587 BP; 36807 A; 38915 C; 38736 G; 36029 T; 100 other;

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